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	PCT/JP00/04414 2000-07-03	
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Arg Phe Ser Gly Ser Ser Ser Gly Ala Asp Arg Tyr Leu Ser Ile Ser 65 70 75 80

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Ala Thr Ile Ser Ser Gly Gly Ser Tyr Thr Tyr Tyr Pro Asp Ser Val 50 55 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn Thr Leu Tyr 65 70 75 80

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Lys Leu Lys Gln Asp Gly Ser His Ser Thr Gly Asp Gly Ile Pro Asp 50 55 60

Arg Phe Ser Gly Ser Ser Ser Gly Ala Glu Arg Tyr Leu Thr Ile Ser 65 70 75 80

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Asp Leu Lys Gln Asp Gly Ser His Ser Thr Gly Asp Gly Ile Pro Asp 50 55 60

Arg Phe Ser Gly Ser Ser Ser Gly Ala Glu Arg Tyr Leu Thr Ile Ser

Ser Leu Gln Ser Glu Asp Glu Ala Asp Tyr Tyr Cys Gly Val Gly Asp 85 90 95

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Asp Leu Lys Gln Asp Gly Ser His Ser Thr Gly Asp Gly Ile Pro Asp 50 55 60

Arg Phe Ser Gly Ser Ser Ser Gly Ala Glu Arg Tyr Leu Thr Ile Ser 65 70 75 80

Ser Leu Gln Ser Glu Asp Glu Ala Asp Tyr Tyr Cys Gly Val Gly Asp 85 90 95

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Ile Glu Trp Tyr Gln Gln Gln Pro Glu Lys Gly Pro Arg Tyr Leu Met 35 40 45

Asp Leu Lys Gln Asp Gly Ser His Ser Thr Gly Asp Gly Ile Pro Asp 50 55 60

Arg Phe Ser Gly Ser Ser Ser Gly Ala Glu Arg Tyr Leu Thr Ile Ser 65 70 75 80

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Asp Leu Lys Gln Asp Gly Ser His Ser Thr Gly Asp Gly Ile Pro Asp 50 55 60

Arg Phe Ser Gly Ser Ser Ser Gly Ala Glu Arg Tyr Leu Thr Ile Ser 65 70 75 80

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Asp Leu Lys Gln Asp Gly Ser His Ser Thr Gly Asp Gly Ile Pro Asp 50 55 60

Arg Phe Ser Gly Ser Ser Ser Gly Ala Glu Arg Tyr Leu Thr Ile Ser

Ser Leu Gln Ser Glu Asp Glu Ala Asp Tyr Ile Cys Gly Val Gly Asp 85 90 95

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Asp Leu Lys Gln Asp Gly Ser His Ser Thr Gly Asp Gly Ile Pro Asp 50 55 60

Arg Phe Ser Gly Ser Ser Ser Gly Ala Glu Arg Tyr Leu Thr Ile Ser 65 70 75 80

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Asp Leu Lys Gln Asp Gly Ser His Ser Thr Gly Asp Gly Ile Pro Asp 50 55 60

Arg Phe Ser Gly Ser Ser Ser Gly Ala Glu Arg Tyr Leu Thr Ile Ser 65 70 75 80

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Asp Leu Lys Gln Asp Gly Ser His Ser Thr Gly Asp Gly Ile Pro Asp 50 55 60

Arg Phe Ser Gly Ser Ser Ser Gly Ala Glu Arg Tyr Leu Thr Ile Ser 65 70 75 80

Ser Leu Gln Ser Glu Asp Glu Ala Asp Tyr Ile Cys Gly Val Gly Asp 85 90 95

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Ala Thr Ile Ser Ser Gly Gly Ser Tyr Thr Tyr Tyr Pro Asp Ser Val 50 55 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr

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Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn 65 70 75	
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tat gtg atg gat ctt aag caa gat gga agc cac agc aca ggt gat ggg Tyr Val Met Asp Leu Lys Gln Asp Gly Ser His Ser Thr Gly Asp Gly 50 55 60	240
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agc att tcc aac atc cag cca gaa gat gaa gca atg tac atc tgt ggt Ser Ile Ser Asn Ile Gln Pro Glu Asp Glu Ala Met Tyr Ile Cys Gly 80 85 90	336
gtg ggt gat aca att aag gaa caa ttt gtg tat gtt ttc ggc ggt ggg Val Gly Asp Thr Ile Lys Glu Gln Phe Val Tyr Val Phe Gly Gly 95 100 105	384
acc aag gtc act gtc cta ggt cag ccc Thr Lys Val Thr Val Leu Gly Gln Pro 110 115	411
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ctg gga gcc Leu Gly Ala 15		_	_			_	_	
acg tac acc Thr Tyr Thr 30	Ile Glu			Gln F			_	.ā
tac ttg atg Tyr Leu Met		-		_	-			_
att cct gat Ile Pro Asp	_		_					
acc atc tcc Thr Ile Ser 80	_	_						
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ctg gga gcc tcg gtc aag ctc acc tgc acc ttg agt agt c Leu Gly Ala Ser Val Lys Leu Thr Cys Thr Leu Ser Ser G 15 20 25	
acg tac acc att gaa tgg tat cag cag cag cca gag aag g Thr Tyr Thr Ile Glu Trp Tyr Gln Gln Gln Pro Glu Lys G 30 35 40	
tac ctg atg gat ctt aag caa gat gga agc cac agc aca g Tyr Leu Met Asp Leu Lys Gln Asp Gly Ser His Ser Thr G 50 55	
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acc atc tcc agc ctc cag tct gag gat gag gct gac tat t Thr Ile Ser Ser Leu Gln Ser Glu Asp Glu Ala Asp Tyr T 80 85 90	
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		tcc Ser -1	_		 _			_		_		_		9	6
_		gcc Ala	_	_										14	4
_		acc Thr		_		-	_	_		-			_	19	2
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		gat Asp	_				_		 _	_	_			28	8
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		gcc Ala	_	_	_						_	_	_		_	144
		acc Thr														192
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tct ttc tcc cag ctt gtg ctg act caa tcg ccc tct gcc tct gcc tcc Ser Phe Ser Gln Leu Val Leu Thr Gln Ser Pro Ser Ala Ser Ala Ser -1 1 5 10	96
ctg gga gcc tcg gtc aag ctc acc tgc acc ttg agt agt cag cac agt Leu Gly Ala Ser Val Lys Leu Thr Cys Thr Leu Ser Ser Gln His Ser 15 20 25	144
acg tac acc att gaa tgg tat cag cag cag cca gag aag ggc cct agg Thr Tyr Thr Ile Glu Trp Tyr Gln Gln Pro Glu Lys Gly Pro Arg 30 35 40 45	192
tac gtg atg gat ctt aag caa gat gga agc cac agc aca ggt gat ggg Tyr Val Met Asp Leu Lys Gln Asp Gly Ser His Ser Thr Gly Asp Gly 50 55 60	240
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gtg ggt gat aca att aag gaa caa ttt gtg tac gtg ttc ggc gga ggg Val Gly Asp Thr Ile Lys Glu Gln Phe Val Tyr Val Phe Gly Gly 95 100 105	384
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tot the tee caq ett qtg etg act caa teg eee tet gee tet gee tee
                                                                       96
Ser Phe Ser Gln Leu Val Leu Thr Gln Ser Pro Ser Ala Ser Ala Ser
        -1 1
ctg gga gcc tcg gtc aag ctc acc tgc acc ttg agt agt cag cac agt
                                                                      144
Leu Gly Ala Ser Val Lys Leu Thr Cys Thr Leu Ser Ser Gln His Ser
    15
                        20
acg tac acc att gaa tgg tat cag cag cag cca gag aag ggc cct aag
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Thr Tyr Thr Ile Glu Trp Tyr Gln Gln Pro Glu Lys Gly Pro Lys
tac ctg atg gat ctt aag caa gat gga agc cac agc aca ggt gat ggg
                                                                      240
Tyr Leu Met Asp Leu Lys Gln Asp Gly Ser His Ser Thr Gly Asp Gly
                50
                                    55
att cet gat ege tte tea gge tee age tet ggg get gag ege tae ete
                                                                      288
Ile Pro Asp Arg Phe Ser Gly Ser Ser Gly Ala Glu Arg Tyr Leu
            65
                                                    75
acc atc tcc agc ctc cag tct gag gat gag gct gac tat atc tgt ggt
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Thr Ile Ser Ser Leu Gln Ser Glu Asp Glu Ala Asp Tyr Ile Cys Gly
                            85
gtg ggt gat aca att aag gaa caa ttt gtg tac gtg ttc ggc gga ggq
                                                                      384
Val Gly Asp Thr Ile Lys Glu Gln Phe Val Tyr Val Phe Gly Gly
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                                     -10
tet tte tee cag ett gtg etg act caa teg eec tet gee tet gee tee
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Ser Phe Ser Gln Leu Val Leu Thr Gln Ser Pro Ser Ala Ser Ala Ser
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ctg gga gcc tcg gtc aag ctc acc tgc acc ttg agt agt cag cac agt
                                                                      144
Leu Gly Ala Ser Val Lys Leu Thr Cys Thr Leu Ser Ser Gln His Ser
    15
                        20
                                             25
acg tac acc att gaa tgg tat cag cag cag cca gag aag ggc cct agg
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Thr Tyr Thr Ile Glu Trp Tyr Gln Gln Pro Glu Lys Gly Pro Arq
30
                    35
                                                             45
tac ctg atg gat ctt aag caa gat gga agc cac agc aca ggt gat ggg
                                                                      240
Tyr Leu Met Asp Leu Lys Gln Asp Gly Ser His Ser Thr Gly Asp Gly
att cct qat cqc ttc tca qqc tcc aqc tct qqq qct qaq cqc tac ctc
                                                                      288
Ile Pro Asp Arg Phe Ser Gly Ser Ser Ser Gly Ala Glu Arg Tyr Leu
            65
acc atc tcc agc ctc cag tct gag gat gag gct gac tat atc tgt ggt
                                                                      336
Thr Ile Ser Ser Leu Gln Ser Glu Asp Glu Ala Asp Tyr Ile Cys Gly
        80
gtg ggt gat aca att aag gaa caa ttt gtg tac gtg ttc ggc gga ggq
                                                                      384
Val Gly Asp Thr Ile Lys Glu Gln Phe Val Tyr Val Phe Gly Gly Gly
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acg ta Thr Ty 30			-			_	_	_			_			_	192
tac gt Tyr Va															240
att cc Ile Pr	_	_					_			_		_			288
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acc aa Thr Ly 110	_		_												411
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Thr Ala

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_		gat Asp	_	_	_	_	_				_	_			_	148
		cgt Arg	-													196
_		cca Pro	_	_		_	_		-	_						244
		tca Ser 75		_				_	_	_			_			292
		tct Ser														340
	_	ccc Pro	_	_												388
aaa	gca	cca	ggt	gag	gtg	gtg	gct	gtg	ccc	tgt	ccg	gac	tac	att	tat	436

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					cct Pro							-			_	532	
					ctc Leu					-	_					580	
					att Ile 190											628	
					gtg Val											676	
_	_	_			atc Ile		_		_		_			_	_	724	
_	_		-		ttc Phe	_	_	_	_						_	772	
					gag Glu											820	
					ccg Pro 270		-		_	_	_					868	
_			_		acc Thr						_	-				916	
					gag Glu											964	
					aag Lys											1012	
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Trp	Gly 330	Leu	Pro	Ala	Val	Phe 335	Val	Ala	Val	Trp	Val 340	Ser	Val	Arg	Ala	
					999 Gly 350											1108
					ccc Pro											1156
					gtc Val				_		_	_				1204
					gac Asp											1252
	_	_			atg Met					_				_		1300
					acc Thr 430											1348
					ctc Leu											1396
			-		tgc Cys				_		_			_		1444
		-			aca Thr			_			_	_	_	_	_	1492
					tat Tyr											1540
					ccc Pro 510											1588
		_			gcc Ala							_	_			1636
cat	gcc	aag	cca	ggg	acc	cca	gcc	ctg	gag	acc	ctc	gag	acc	aca	cca	1684

His Ala Ly	s Pro Gly Th 540	r Pro Ala Leu 545	ı Glu Thr Leu Glu	Thr Thr Pro 550
_	t Ala Ala Pr		ggg ttc ctc aac Gly Phe Leu Asn 565	
			g cct gag cgg cca / Pro Glu Arg Pro 580	• •
		g aca gtc atg u Thr Val Met O	-	1810